

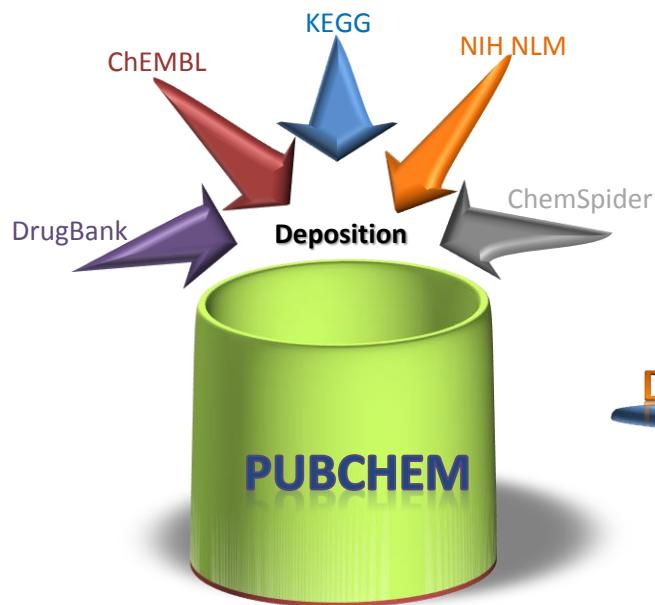
Semantic Annotation of PubChem Database

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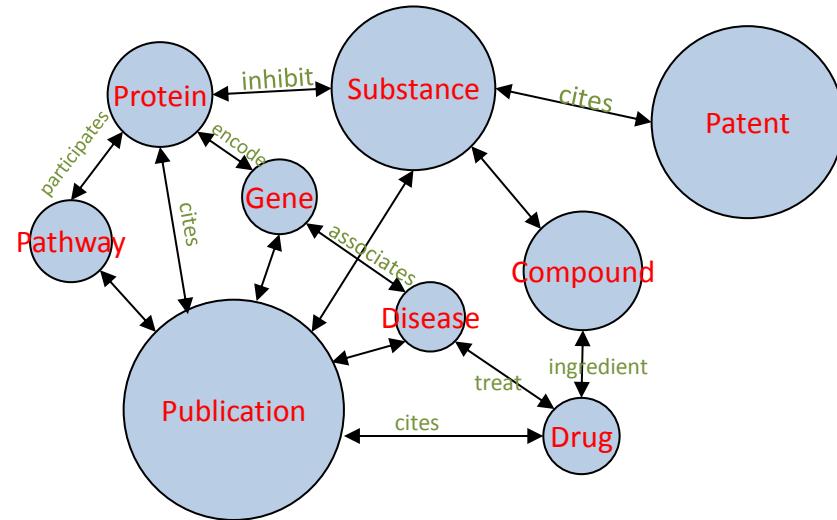
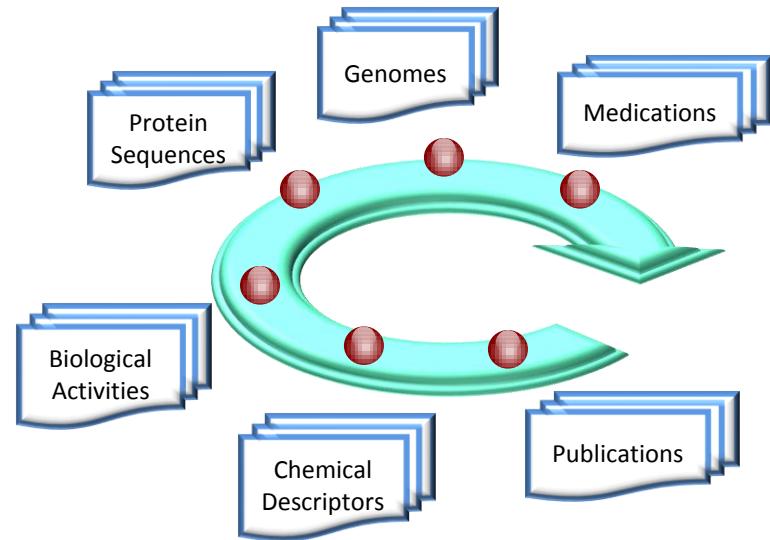


- Background Information
- Ontology-based Data Integration
- PubChem RDF URI References
- PubChem RDF Subdomains
- REST Interface
- Use Cases

Background Information



Data Integration



Semantic Annotation

Standardized Ontologies		
Prefix	Namespace	Vocabularies
rdfs	http://www.w3.org/2000/01/rdf-schema#	RDF Schema
rdf	http://www.w3.org/1999/02/22-rdf-syntax-ns#	RDF
owl	http://www.w3.org/2002/07/owl#	OWL
xsd	http://www.w3.org/2001/XMLSchema#	XML Schema
chebi	http://purl.obolibrary.org/obo/	ChEBI and
uo		UO
sio		CHEMINF and
cheminf	http://semanticscience.org/resource/	SIO
skos	http://www.w3.org/2004/02/skos/core#	SKOS
obo	http://purl.obolibrary.org/obo/	BFO and OBI and IAO
bao	http://www.bioassayontology.org/bao#	BAO
qudt	http://data.nasa.gov/qudt/owl/qudt#	QUDT
cito	http://purl.org/spar/cito/	CiTO
fabio	http://purl.org/spar/fabio/	FaBio
ops	http://www.openphacts.org/units/	Open PHACTS vocabulary
pr	http://purl.obolibrary.org/obo/pr#	PRO
go	http://purl.obolibrary.org/obo/go#	GO
dcterms	http://purl.org/dc/terms/	DCMI Terms
pav	http://purl.org/pav/	PAV
foaf	http://xmlns.com/foaf/0.1/	FOAF vocabulary

PubChemRDF Subdomains

<i>Prefix</i>	<i>Namespace</i>
compound	http://rdf.ncbi.nlm.nih.gov/pubchem/compound/
substance	http://rdf.ncbi.nlm.nih.gov/pubchem/substance/
descr	http://rdf.ncbi.nlm.nih.gov/pubchem/descriptor/
syno	http://rdf.ncbi.nlm.nih.gov/pubchem/synonym/
bioassay	http://rdf.ncbi.nlm.nih.gov/pubchem/bioassay/
measuregroup	http://rdf.ncbi.nlm.nih.gov/pubchem/measuregroup/
endpoint	http://rdf.ncbi.nlm.nih.gov/pubchem/endpoint/
protein	http://rdf.ncbi.nlm.nih.gov/pubchem/protein/
domain	http://rdf.ncbi.nlm.nih.gov/pubchem/domain/
biosystem	http://rdf.ncbi.nlm.nih.gov/pubchem/biosystem/
gene	http://rdf.ncbi.nlm.nih.gov/pubchem/gene/
reference	http://rdf.ncbi.nlm.nih.gov/pubchem/reference/
nbr	http://rdf.ncbi.nlm.nih.gov/pubchem/neighbor/
source	http://rdf.ncbi.nlm.nih.gov/pubchem/source/
vocab	http://rdf.ncbi.nlm.nih.gov/pubchem/vocabulary#

<http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID60823>

<http://rdf.ncbi.nlm.nih.gov/pubchem/substance/SID103554720>

http://rdf.ncbi.nlm.nih.gov/pubchem/descriptor/CID60823_Molecular_Weight

http://rdf.ncbi.nlm.nih.gov/pubchem/synonym/MD5_9a05646d461669f86de312d88ab5748a

<http://rdf.ncbi.nlm.nih.gov/pubchem/bioassay/AID1788>

<http://rdf.ncbi.nlm.nih.gov/pubchem/measuregroup/AID447528>

http://rdf.ncbi.nlm.nih.gov/pubchem/measuregroup/AID1788_1

http://rdf.ncbi.nlm.nih.gov/pubchem/measuregroup/AID363_PMID16161995

http://rdf.ncbi.nlm.nih.gov/pubchem/endpoint/SID103164874_AID443491

http://rdf.ncbi.nlm.nih.gov/pubchem/endpoint/SID99445338_AID2202_1

http://rdf.ncbi.nlm.nih.gov/pubchem/endpoint/SID8033500_AID363_PMID10395478

<http://rdf.ncbi.nlm.nih.gov/pubchem/protein/GI124375976>

<http://rdf.ncbi.nlm.nih.gov/pubchem/reference/PMID10395478>

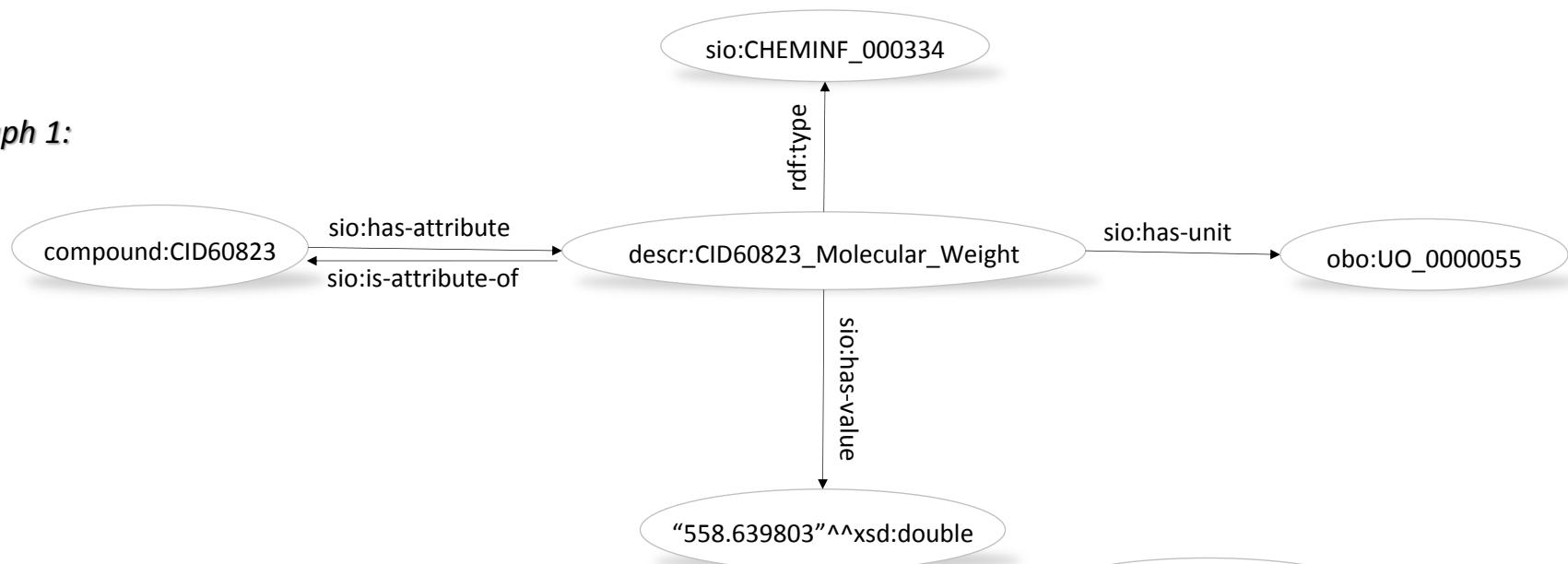
<http://rdf.ncbi.nlm.nih.gov/pubchem/source/ChEMBL>

http://rdf.ncbi.nlm.nih.gov/pubchem/neighbor/CID60823_CID68019409_2DSimilarity

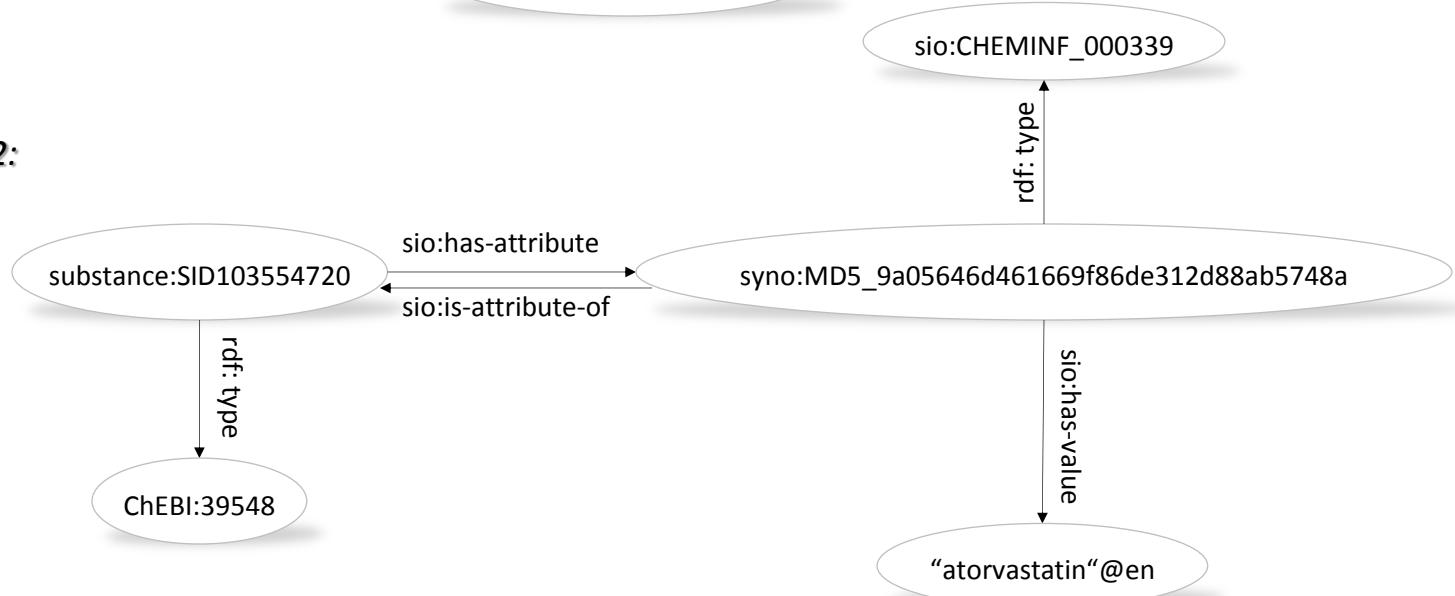
http://rdf.ncbi.nlm.nih.gov/pubchem/neighbor/CID60823_CID11330946_3DSimilarity

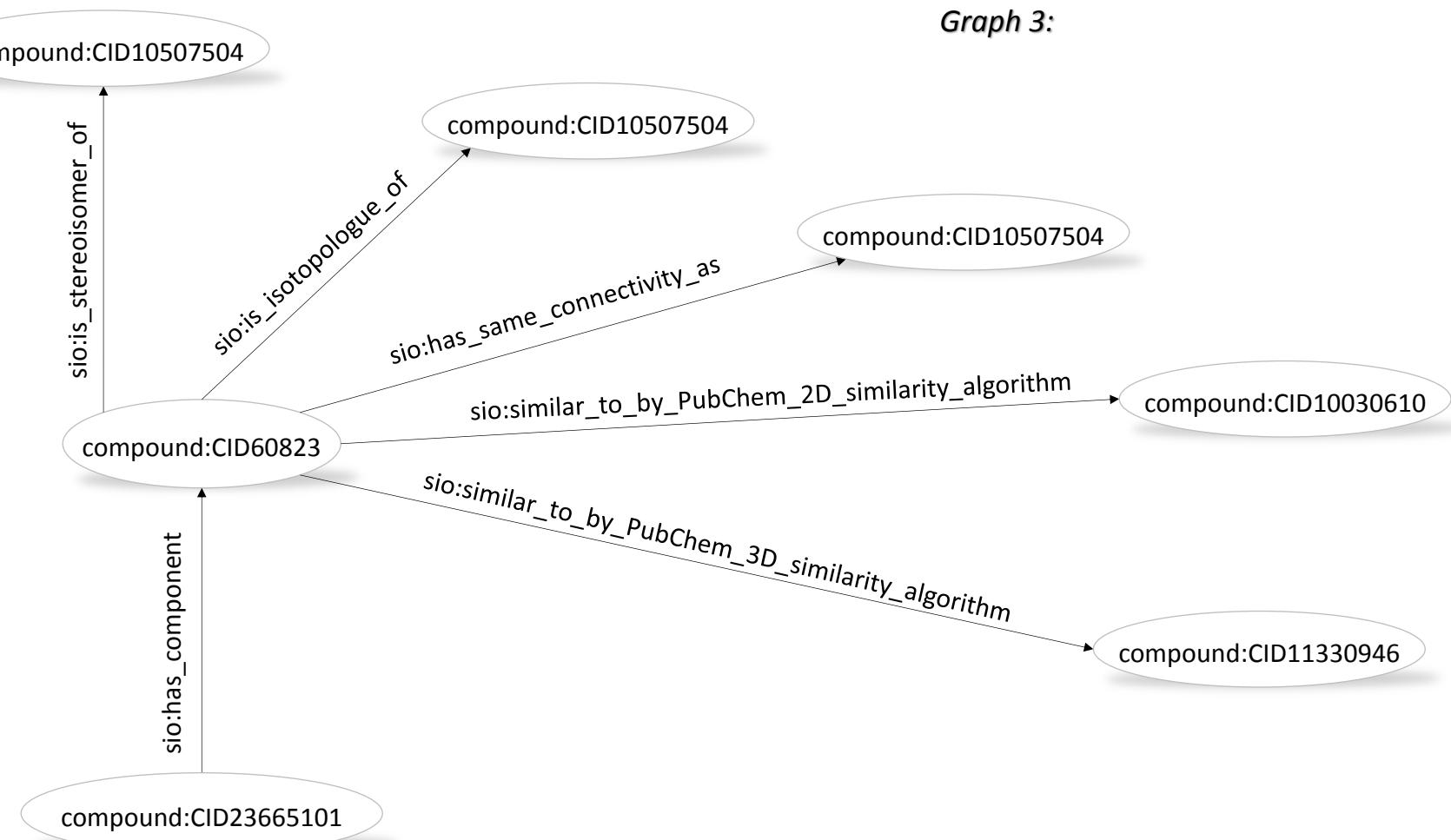
http://rdf.ncbi.nlm.nih.gov/pubchem/neighbor/GI548481_GI129900_SequenceSimilarity

Graph 1:

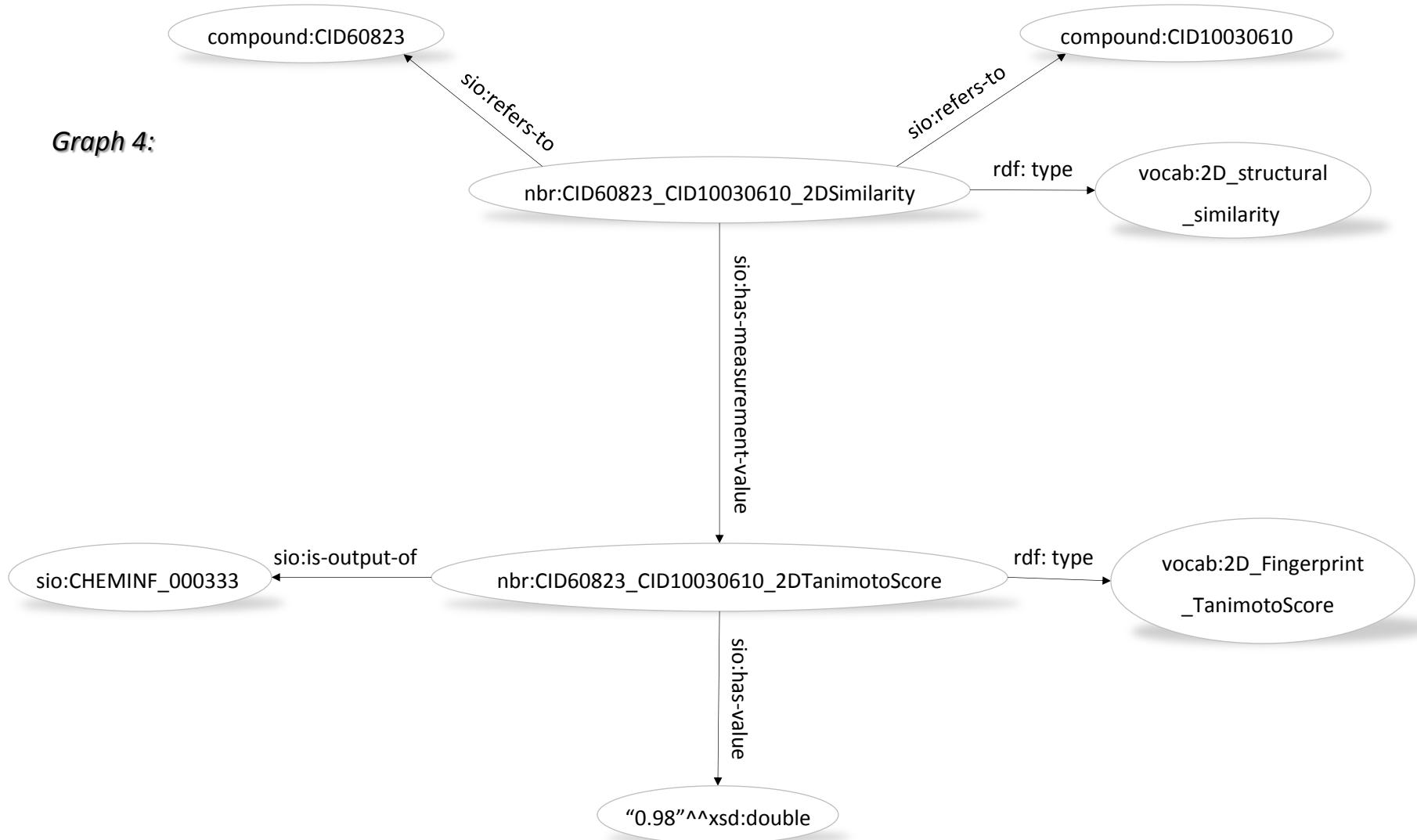


Graph 2:

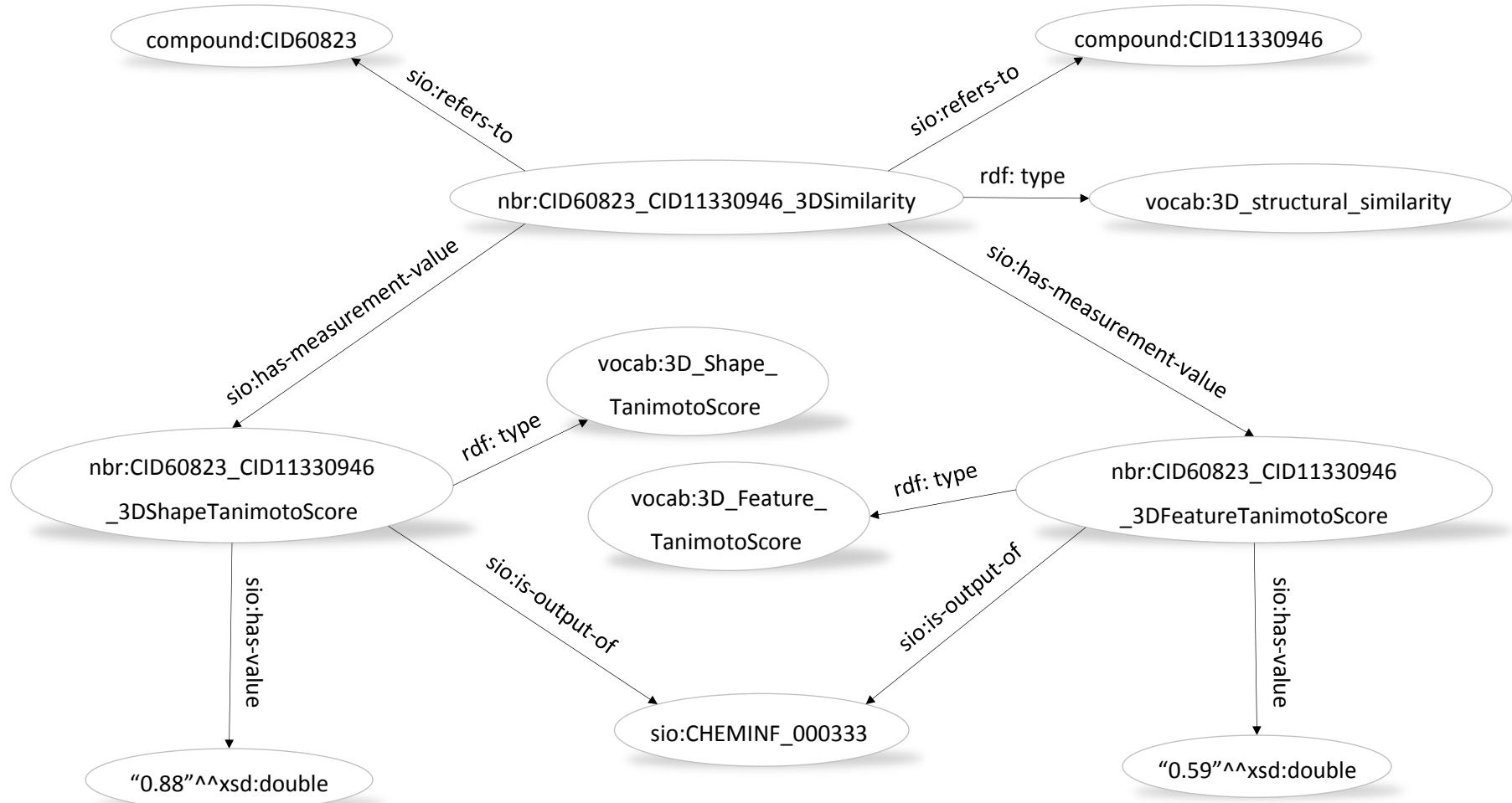




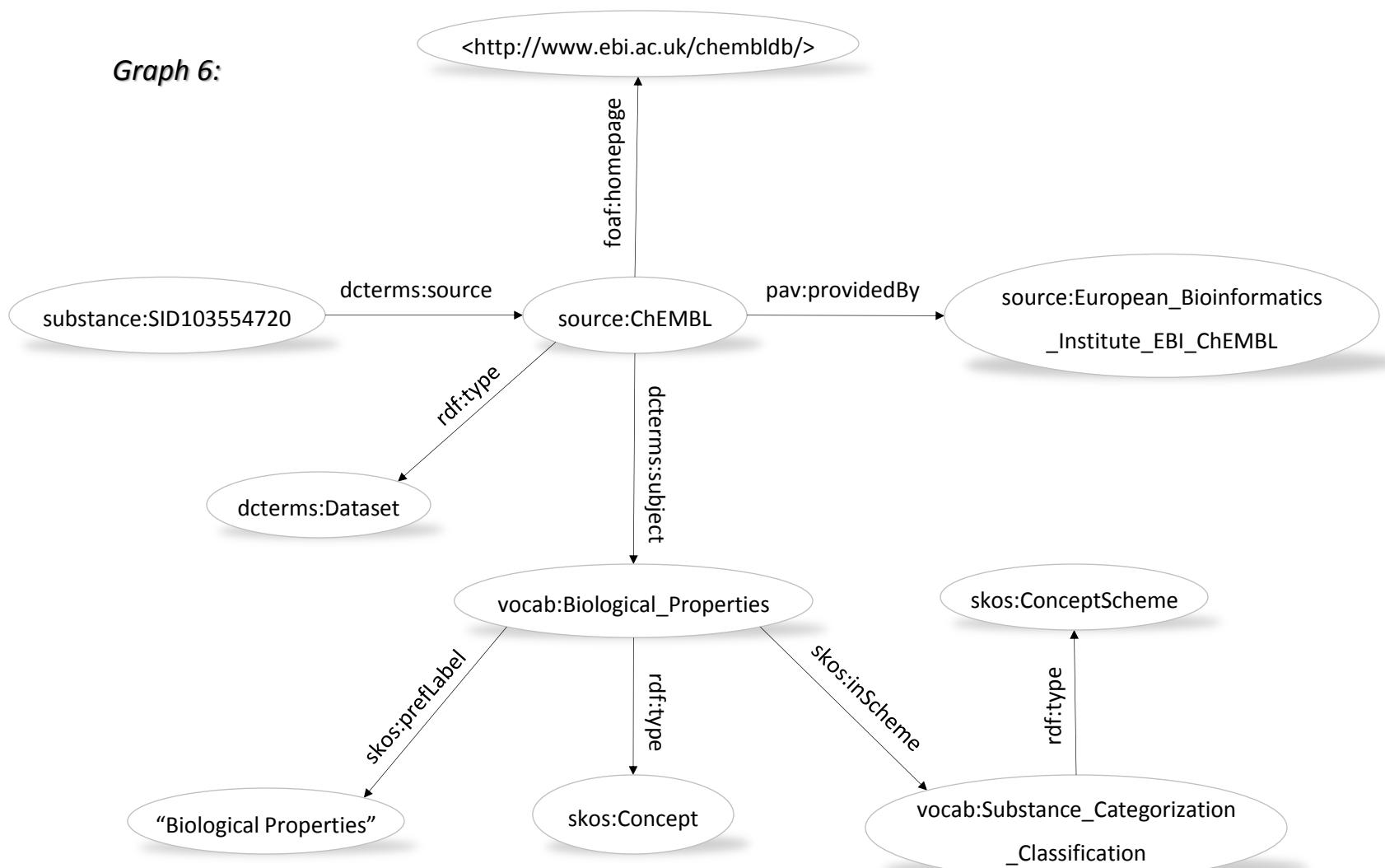
Graph 4:



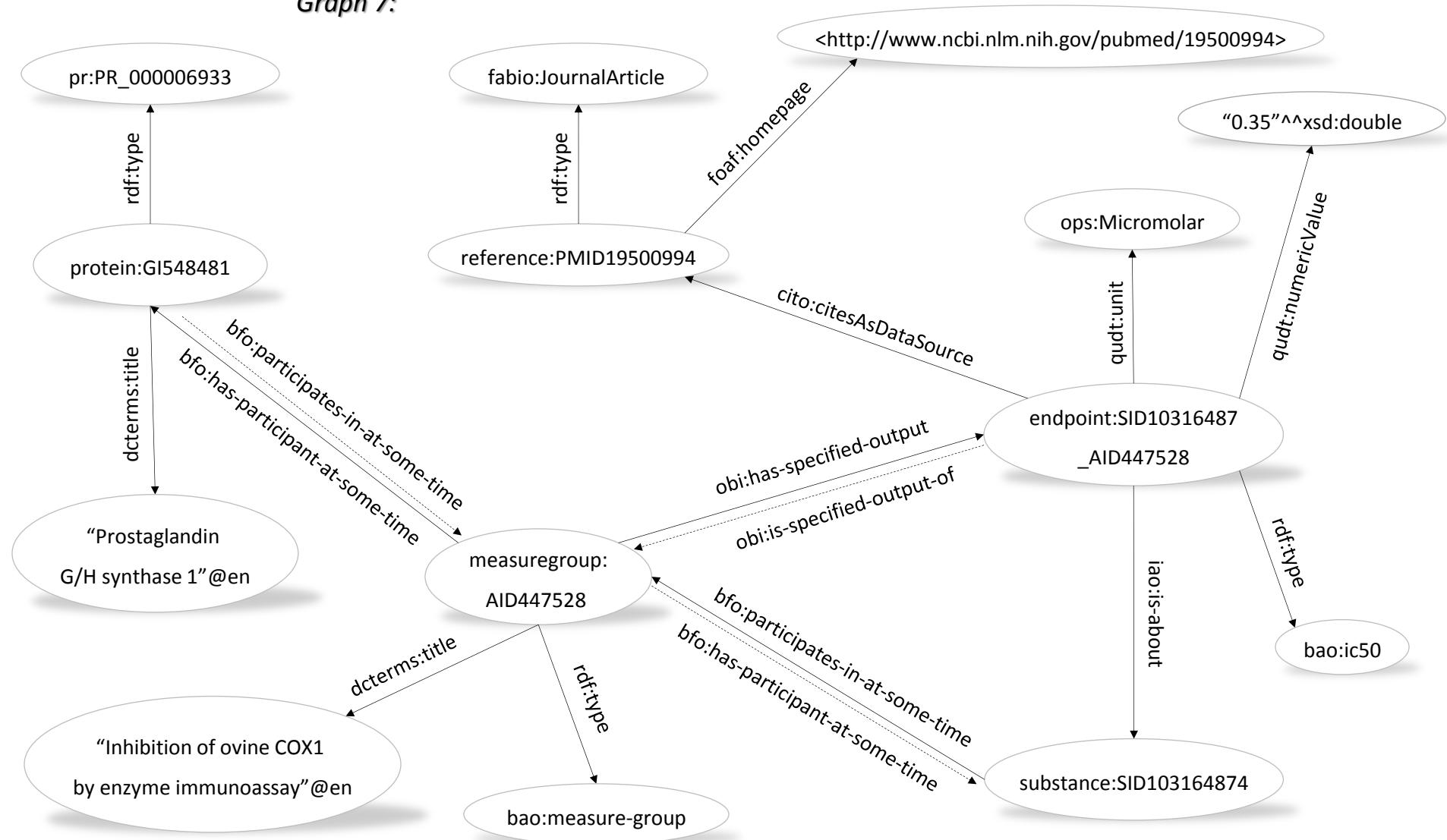
Graph 5:



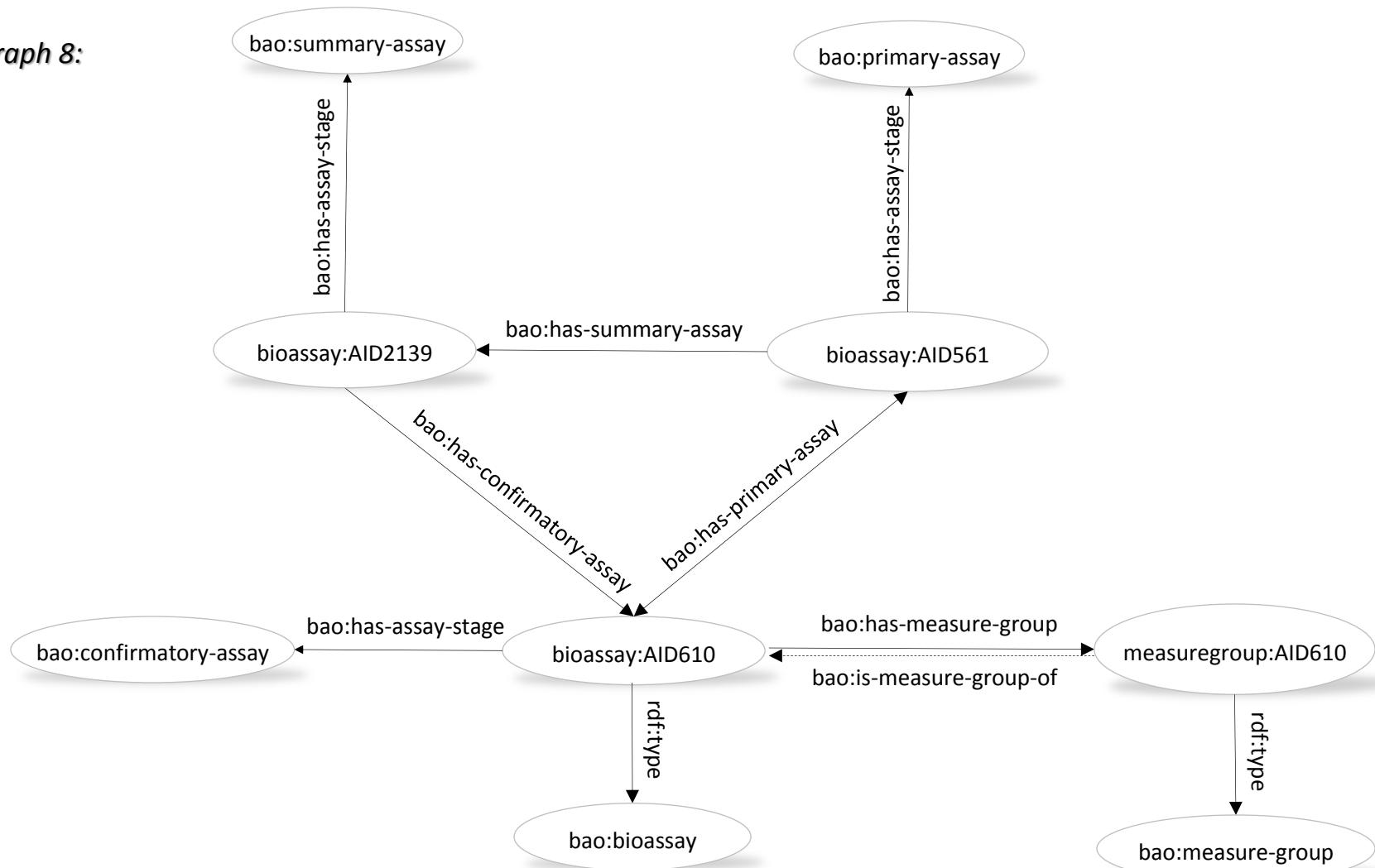
Graph 6:



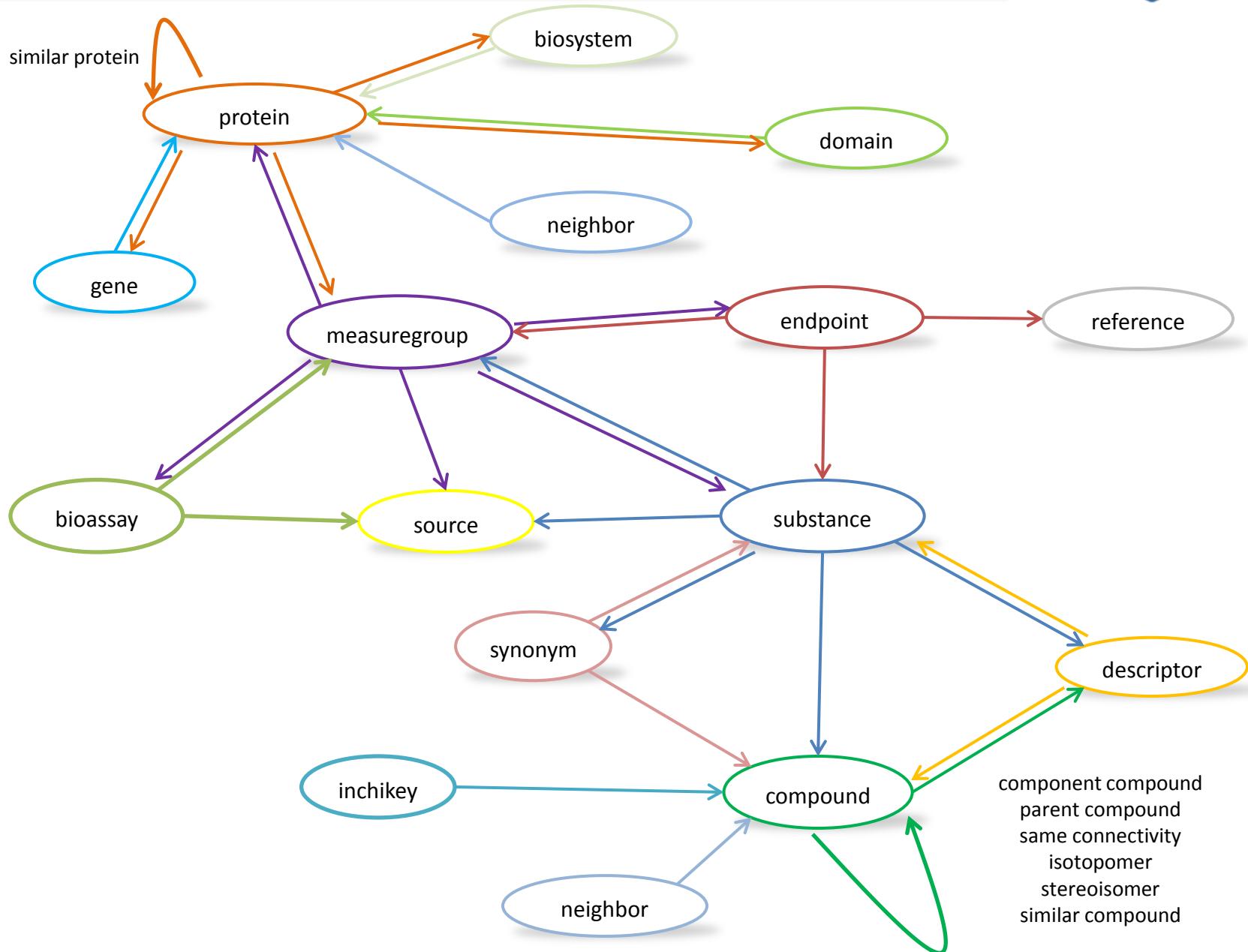
Graph 7:



Graph 8:



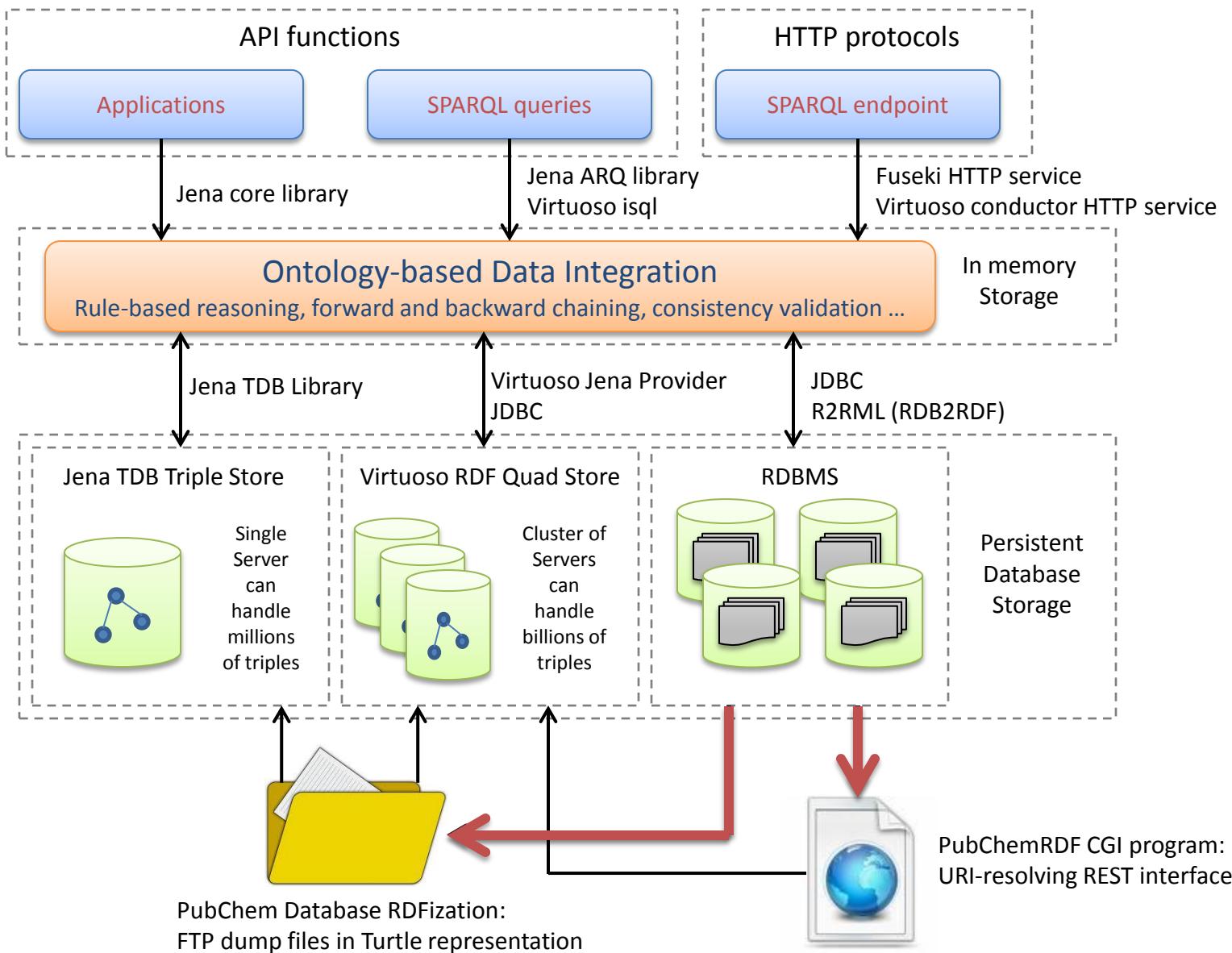
PubChem RDF Linkage between Subdomains



- <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244.rdf>
- <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244.xml>
- <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244.rdfxml>
- <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244.html>
- <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244.turtle>
- <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244.ttl>
- <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244.json>
- <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244.ntriples>

HTTP Status	Description
400	Bad Query URL or Request URI
404	Input URI is invalid or cannot be identified in databases
405	MIME output format is unspecified or invalid
500	Some problem on the server side occurs
504	The request timed out (over 28 second)

MIME Type	HTTP Accept Header or URI extension
application/rdf+xml+abb rev	default application/rdf+xml+abbrev rdfxml-abbrev
application/rdf+xml	application/rdf+xml text/rdf rdfxml rdf xml
application/xhtml+xml	application/xhtml+xml text/html html
application/x-turtle^a	application/rdf+n3 application/turtle application/x-turtle turtle ttl
application/json^b	application/json text/json json
text/plain	text/plain ntriples
text/rdf+n3	text/n3 text/rdf+n3 Accept: n3



Java - jena_test/src/example/Jena_pc_test.java - Eclipse

File Edit Source Refactor Navigate Search Project Run Window Help

Package Explorer JenaFunctions.java JenaTest.java Jena_pc_test.java

```

1 package example;
2
3 import java.util.Iterator;
4
5 import org.apache.log4j.PropertyConfigurator;
6
7 import com.hp.hpl.jena.query.Dataset;
8 import com.hp.hpl.jena.query.ReadWrite;
9 import com.hp.hpl.jena.sparql.core.Quad;
10 import com.hp.hpl.jena.tdb.TDBFactory;
11 import com.hp.hpl.jena.tdb.TDBLoader;
12 import com.hp.hpl.jena.tdb.sys.TDBInternal;
13
14 public class Jena_pc_test {
15
16
17     public static void main(String[] args) {
18
19         PropertyConfigurator.configure("log4j.properties");
20
21         String dataSet = "target/pc_comp";
22         String filename1 = "data/pc_comp.ttl";
23         String filename2 = "data/pc_comp_descr.ttl";
24
25         TDBLoader.load( TDBInternal.getBaseDatasetGraphTDB( TDBFactory.createDatasetGraph( dataSet ) ), filename1, true );
26         TDBLoader.load( TDBInternal.getBaseDatasetGraphTDB( TDBFactory.createDatasetGraph( dataSet ) ), filename2, true );
27
28         Dataset dataset = TDBFactory.createDataset( dataSet );
29         dataset.begin(ReadWrite.READ);
30         try {
31             Iterator<Quad> iter = dataset.asDatasetGraph();
32             int count = 0;
33             while ( iter.hasNext() ) {
34                 Quad quad = iter.next();
35                 System.out.println(quad);
36                 count++;
37             }
38             System.out.println( count );
39         } finally {
34         dataset.end();
35     }
36 }
37
38
39
40
41
42
43 }
44

```

Jena libraries:

- Core
- ARQ
- IRI
- TDB

TDBLoader.load() function:
Bulk load RDF triples from files

Problems @ Javadoc Declaration Console Progress

<terminated> Jena_pc_test [Java Application] C:\Program Files\Java\jre7\bin\javaw.exe (Sep 4, 2013 2:33:28 PM)

[urn:x-arq:DefaultGraph http://rdf.ncbi.nlm.nih.gov/pubchem/chemical-descriptor/CID99999_Undefined_Bond_Stereo_Count http://semanticscience.org/resource/10440629]

PubChem RDF Utility



Java - virtuoso_test/src/example/virtuosoTest.java - Eclipse

File Edit Source Refactor Navigate Search Project Run Window Help

Package Explorer

- db_test
- jena_test
- paxtool_test
- pdfbox_test
- virtuoso_test
 - src
 - example
 - virtuosoTest.java
- JRE System Library [JavaSE-1.7]
- virtuoso
 - arq.jar - C:\Users\fug2\Documents\Apache-Jena\virtuoso-jena
 - axis.jar - C:\Users\fug2\Documents\Apache-Jena\virtuoso-jena
 - commons-logging.jar - C:\Users\fug2\Documents\Apache-Jena\virtuoso-jena
 - icu4j_3_4.jar - C:\Users\fug2\Documents\Apache-Jena\virtuoso-jena
 - iri.jar - C:\Users\fug2\Documents\Apache-Jena\virtuoso-jena
 - jena.jar - C:\Users\fug2\Documents\Apache-Jena\virtuoso-jena
 - virt_jena.jar - C:\Users\fug2\Documents\Apache-Jena\virtuoso-jena
 - virtjdbc3.jar - C:\Users\fug2\Documents\Apache-Jena\virtuoso-jena
 - xercesimpl.jar - C:\Users\fug2\Documents\Apache-Jena\virtuoso-jena

Graph.read() function
REST Interface: <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244>
MIME Type: RDF/XML

```
public static void main(String[] args) {
    String url;
    if(args.length == 0)
        url = "jdbc:virtuoso://lmem04:1111";
    else
        url = args[0];
    /* STEP 1 */
    VirtGraph graph = new VirtGraph ("http://PubChemRDF/CID2244", url, "dba", "PubChemRDF");
    /* STEP 2 */
    /* Load data */
    graph.clear ();
    System.out.print ("Begin read from 'http://pubchem.ncbi.nlm.nih.gov/rest/rdf/compound/CID2244.html'. \n");
    graph.read("http://pubchem.ncbi.nlm.nih.gov/rest/rdf/compound/CID2244", "RDF/XML");
    System.out.println ("Done. \n");
    /* STEP 3 */
    while (results.hasNext()) {
        QuerySolution result = results.nextSolution();
        RDFNode graph_name = result.get("graph");
        RDFNode s = result.get("s");
        RDFNode p = result.get("p");
        RDFNode o = result.get("o");
        System.out.println(graph_name + " { " + s + " " + p + " " + o + " . }");
    }
    System.out.println("graph.getCount() = " + graph.getCount());
}
```

Problems @ Javadoc Declaration Console Progress

<terminated> virtuosoTest (1) [Java Application] C:\Program Files\Java\jre7\bin\javaw.exe (Sep 3, 2013 11:21:14 AM)
http://PubChemRDF/CID2244 { http://rdf.ncbi.nlm.nih.gov/pubchem/compound/CID2244 http://semanticsscience.org/resource/has-attribute ht
graph.getCount() = 11100

Jena Framework
Virtuoso Jena
Provider
JDBC Driver

Jena libraries:

- Core
- ARQ
- IRI
- TDB

The screenshot shows an Eclipse IDE interface with the following details:

- Project Explorer:** Shows the project structure under "jena_test". It includes a "src" folder containing "example" (with "BiosystemsRDF.java", "BiosystemsTest.java", "Jena_pc_test.java", "JenaFunctions.java", and "JenaTest.java"), and JRE System Library [JavaSE-1.6] and Jena dependencies.
- Code Editor:** Displays the content of "Jena_pc_test.java". The code uses Jena's Query API to construct and execute a SPARQL query against a dataset. A red box highlights the query string, and another red box highlights the "QueryFactory.create(queryString)" and "QueryExecution qexec = QueryExecutionFactory.create(query, dataset);".
- Annotations:**
 - A red box encloses the SPARQL query string, which is used to filter compounds based on their MW, HD, HA, and LogP values.
 - A red box encloses the "QueryFactory.create(queryString)" and "QueryExecution qexec = QueryExecutionFactory.create(query, dataset);" lines, indicating the functions used to create and execute the query.
 - A red arrow points from the "Jena libraries:" box to the Jena dependencies in the Project Explorer.
 - A red arrow points from the "QueryFactory and QueryExecution functions" box to the highlighted code in the editor.
- Bottom Status Bar:** Shows "Jena_pc_test [Java Application]" and the command line "C:\Program Files\Java\jre7\bin\javaw.exe (Sep 4, 2013 3:40:18 PM)".

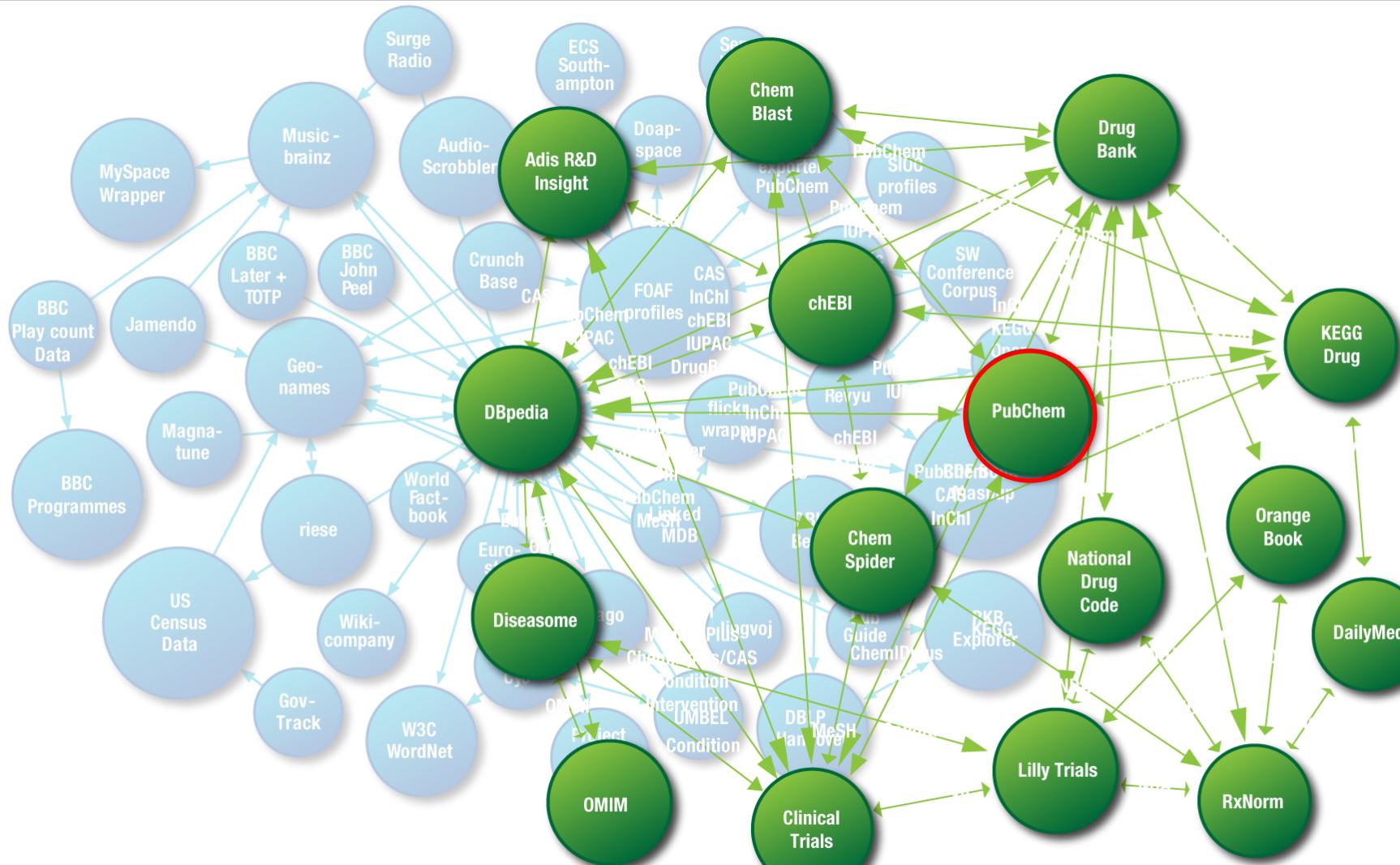
```

1 package example;
2
3 import org.apache.log4j.PropertyConfigurator;
4
5 public class Jena_pc_test {
6
7     public static void main(String[] args) {
8
9         PropertyConfigurator.configure("log4j.properties");
10
11         String dataSet = "target/pc_comp";
12         Dataset dataset = TDBFactory.createDataset( dataSet );
13
14
15         String queryString =
16             "PREFIX compound: <http://rdf.ncbi.nlm.nih.gov/pubchem/compound/> " +
17             "PREFIX rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>" +
18             "PREFIX sio: <http://semanticscience.org/resource/>" +
19             "SELECT DISTINCT ?compound " +
20             "WHERE ? " +
21             " ?compound sio:has-attribute ?MW . " +
22             " ?MW rdf:type sio:CHEMINF_000334 ; " +
23             " ?sio:has-value ?MWValue . " +
24             " ?compound sio:has-attribute ?HD . " +
25             " ?HD rdf:type sio:CHEMINF_000387 ; " +
26             " ?sio:has-value ?HDCount . " +
27             " ?compound sio:has-attribute ?HA . " +
28             " ?HA rdf:type sio:CHEMINF_000388 ; " +
29             " ?sio:has-value ?HACount . " +
30             " ?compound sio:has-attribute ?LogP . " +
31             " ?LogP rdf:type sio:CHEMINF_000395 ; " +
32             " ?sio:has-value ?LPValue . " +
33             " FILTER ( ?MWValue < 500 && ?HDCount < 5 && ?HACount < 10 && ?LPValue < 5 ) " +
34             " } ";
35
36         Query query = QueryFactory.create(queryString);
37
38         QueryExecution qexec = QueryExecutionFactory.create(query, dataset);
39
40
41         try
42         {
43             ResultSet results = qexec.execSelect();
44             ResultSetFormatter.out( results, query )
45         } finally {
46             qexec.close();
47         }
48
49     }
50
51 }
52
53
54
55
56
57
58

```

- PubChem RDF is intended for ontology-based data integration
- PubChem databases have been semantically exposed to linked open data
- REST interface can be accessed to resolve URI references
- FTP dump files can be bulk-loaded into open source triples stores

Conclusion



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Uma Vempati
Egon Willighagen

Thank you and Questions!